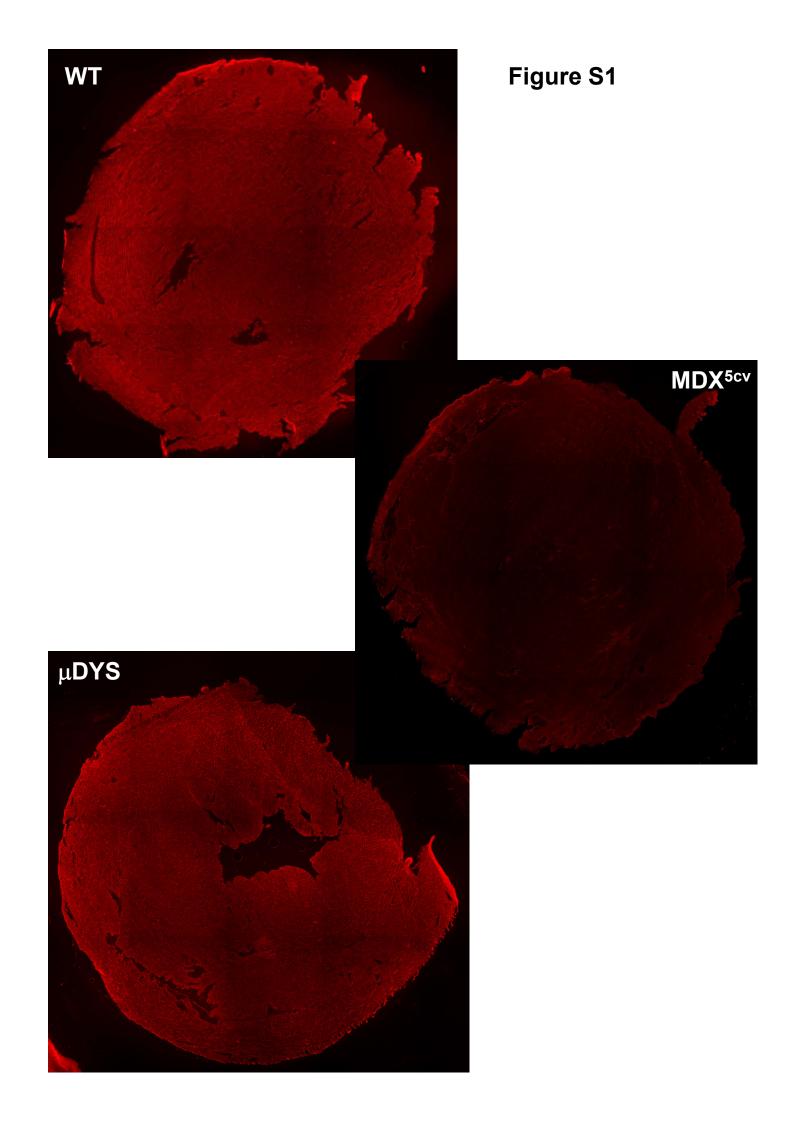
Figure S1:  $\mu$ DYS is uniformly expressed in the myocardium of transgenic  $\mu$ DYS- $mdx^{5cv}$  mice up to 1 year of age. Montages of cardiac sections from 1-year old wild type (WT),  $mdx^{5cv}$  (MDX) and  $\mu$ DYS- $mdx^{5cv}$  ( $\mu$ DYS) mice stained with the MANEX1011B antibody (red) that recognizes an epitope present in both full length and micro-dystrophin.



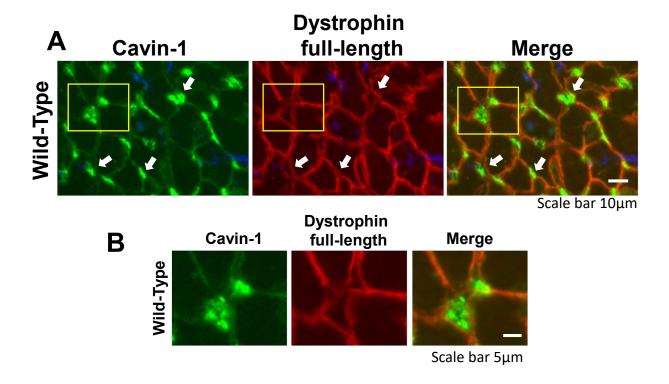
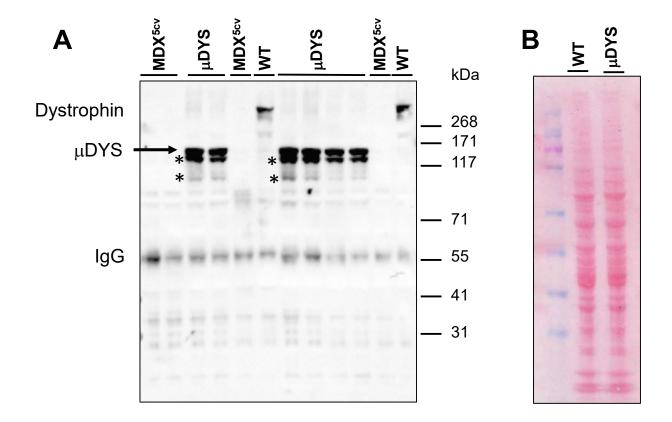


Figure S2: Full length dystrophin is expressed in cardiomyocytes and is absent from capillaries. A. Representative images of cardiac sections from a 6 months old wild-type male mouse double-labelled with the MANEX1011B antibody (red) that only recognizes full-length dystrophin, and with an antibody to cavin-1 (green) a protein highly expressed in capillaries in the mouse myocardium. Scale bar =  $10\mu m$ . B. Higher magnification of the boxed area in A to highlight the lack of dystrophin (red) expression in capillaries that are highly positive for cavin-1 (green) and in the interstitial tissue in-between cardiomyocytes. Note that cavin-1 is also expressed at lower levels at the membrane of cardiomyocytes where it co-localizes with dystrophin. Scale bar =  $5\mu m$ .



**Figure S3:** μDYS stability *in vivo*. **A.** Full nitrocellulose membrane of cardiac lysates from wild type (WT),  $mdx^{5cv}$  (MDX) and μDYS- $mdx^{5cv}$  (μDYS) mice probed with the MANEX1011B antibody. The image was over-exposed to reveal the presence of additional lower bands (asterisks) and a smear below the 144 kDa μDys band (arrow) in the μDYS- $mdx^{5cv}$  samples. **B.** Representative nitrocellulose membrane stained with Ponceau S to visualize all proteins in heart lysates from a wild-type and a μDYS- $mdx^{5cv}$  mouse. Both samples show well-defined protein bands and no smears indicating a lack of global protein degradation.

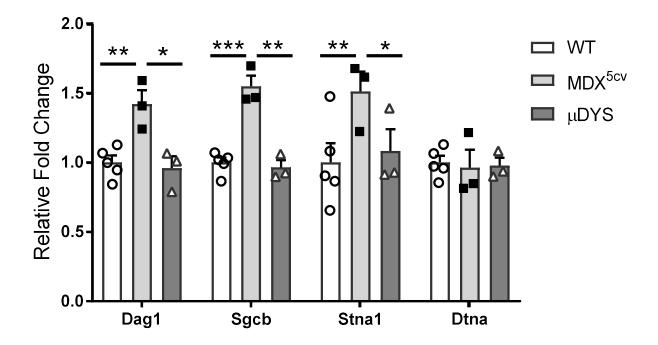


Figure S4: Quantification of DAPC mRNA transcripts in hearts from wild-type (WT),  $mdx^{5cv}$  and  $\mu$ DYS- $mdx^{5cv}$  ( $\mu$ DYS) mice. Transcript amounts for the indicated genes were quantified by quantitative RT-PCR using the ΔΔCt method. Transcript levels were normalized to HPRT and then further normalized to levels in wild-type samples. Statistical differences were determined using an ordinary two-way ANOVA followed by the Tuckey test with correction for multiple comparisons. \* p<0.05; \*\* p<0.01; \*\*\* p<0.005. DAG1: dystroglycan; SGCB: β-sarcoglycan; STNA1:  $\alpha$ 1-syntrophin; DTNA:  $\alpha$ -dystrobrevin.

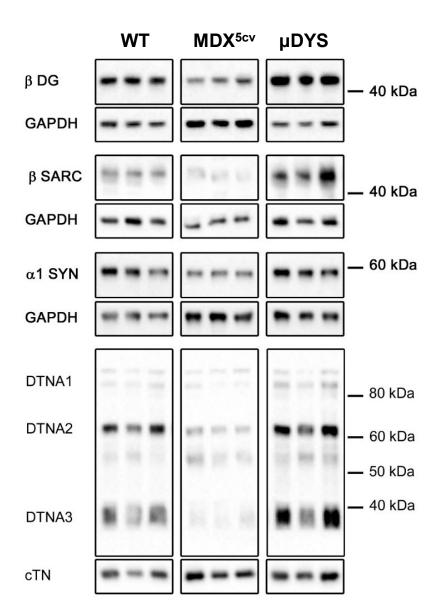


Figure S5: Representative western blots of total cardiac lysates from wild-type (WT), *mdx*<sup>5cv</sup> and μDYS-*mdx*<sup>5cv</sup> (μDYS) mice probed for DAPC proteins. After transfer, nitrocellulose membranes were cut at appropriate molecular weights to allow probing for multiple proteins, including loading controls (GAPDH or cardiac troponin (cTN) depending on the protein molecular weight) on the same gel. Each membrane had 3-4 samples corresponding to individual mice for each genotype. Densitometry was performed using a Chemidoc apparatus and intensity of bands for the proteins of interest was first normalized to the loading control on that exact same sample, and then to the reference wild-type samples on the same membrane.

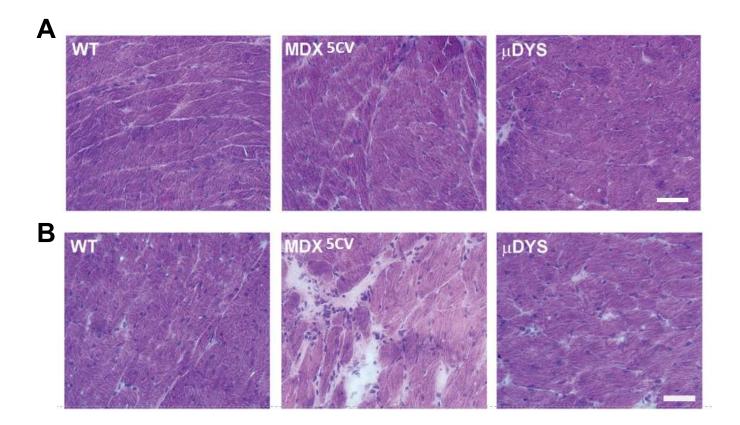


Figure S6:  $\mu$ DYS expression rescues cardiac histopathology in  $mdx^{5cv}$  mice. Representative pictures of Hematoxylin and Eosin stained heart sections from wild type (WT),  $mdx^{5cv}$  and  $\mu$ DYS- $mdx^{5cv}$  mice ( $\mu$ DYS) at 6 months (**A**) and 1 year of age (**B**). Scale bars: 50 $\mu$ m.

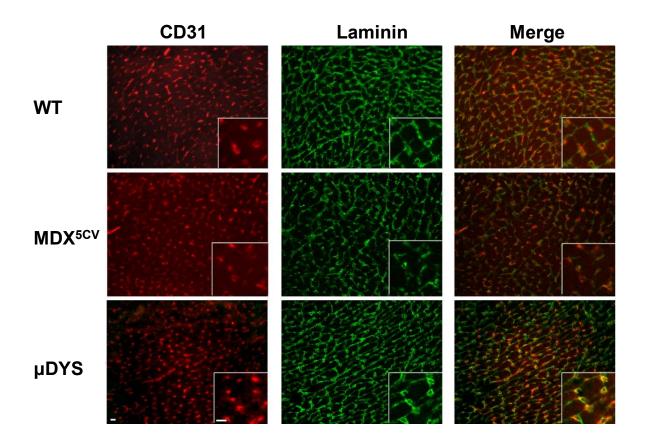


Figure S7: Immuno-labeling for quantification of capillary density and cardiomyocyte size. Representative pictures of heart sections from wild type (WT), *mdx*<sup>5cv</sup> (MDX) and μDYS-*mdx*<sup>5cv</sup> (μDYS) mice immunolabelled for CD31 (red) to visualize capillaries and a pan-laminin (green) to visualize boundaries of cardiomyocytes and blood vessels. Insets show magnifications to highlight capillaries recognizable by the expression of CD31 within a thin ring of laminin staining that is less than 10μm in diameter. Pictures shown are for mice at 6 months of age. Scale bars: 20μm. Uneven brightness of staining is due to the use of Sudan black to quench tissue auto-fluorescence. Images have been processed in Photoshop to increase contrast at the edges of the field and create a more even view of the staining. However, quantifications shown in Figure 1 were performed on unprocessed images.

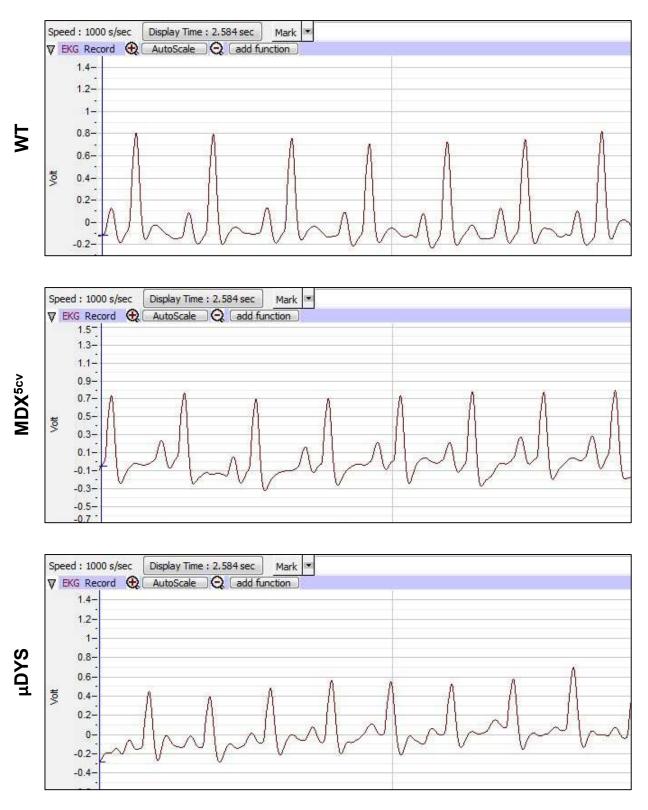


Figure S8: Representative electrocardiogram traces from wild-type,  $mdx^{5cv}$  and  $\mu DYS-mdx^{5cv}$  mice.

Α					
1	MEDVTLHIVE	R <b>PYSGFPDAS</b>	SEGPEPTQGE	ARATEEPSGT	GSDELIKSDQ
51	VNGVLVLSLL	DKIIGAVDQI	QLTQAQLEER	QAEMEGAVQS	<b>iqgelsk</b> lgk
101	AHATTSNTVS	<b>K</b> LLEKVRKVS	VNVKTVRGSL	ERQAGQIK <b>KL</b>	<b>EVNEAELLR</b> R
151	RNFK <b>VMIYQD</b>	<b>EVK</b> LPAKLSV	SKSLKESEAL	PEKEGDELGE	GERPEDDTAA
201	IELSSDEAVE	VEEVIEESRA	ERIKRSGLRR	VDDFKKAFSK	EKMEKTKVRT
251	RENLEKTRLK	TKENLEKTRH	TLEKRMNKLG	TRLVPVERRE	KLKTSRDKLR
301	KSFTPDHVVY	<b>ar</b> sktavyk <b>v</b>	PPFTFHVKKI	REGEVEVLKA	TEMVEVGPED
351	DEVGAERGEA	<b>TDLLR</b> GSSPD	VHTLLEITEE	SDAVLVDKSD	SD
В					
1	MEDVTLHIVE	RPYSGFPDAS	SEGPEPTQGE	ARATEEPSGT	GSDELIK <mark>SDQ</mark>
51	VNGVLVLSLL	DKIIGAVDQI	QLTQAQLEER	QAEMEGAVQS	<b>iqgelsk</b> lgk
101	AHATTSNTVS	KLLEKVRKVS	VNVKTVRGSL	ERQAGQIKKL	EVNEAELLRR
151	RNFKVMIYQD	EVKLPAKLSV	SKSLKESEAL	PEKEGDELGE	GERPEDDTAA
201	IELSSDEAVE	VEEVIEESRA	ERIKRSGLRR	VDDFKKAFSK	EKMEKTKVRT
251	RENLEKTRLK	TKENLEKTRH	TLEKRMNKLG	TRLVPVERRE	KLKTSRDKLR
301	KSFTPDHVVY	ARSKTAVYKV	PPFTFHVKKI	REGEVEVLKA	TEMVEVGPED
351	DEVGAERGEA	TDLLRGSSPD	VHTLLEITEE	SDAVLVDKSD	SD

Figure S9: Cavin-1 peptides identified by proteomics. The full amino acid sequence coverage map of cavin-1 is shown. Amino acids in red correspond to peptides identified by proteomics in dystrophin immunoprecipitations from wild-type (A) and  $\mu$ DYS- $mdx^{5cv}$  (B) mice.

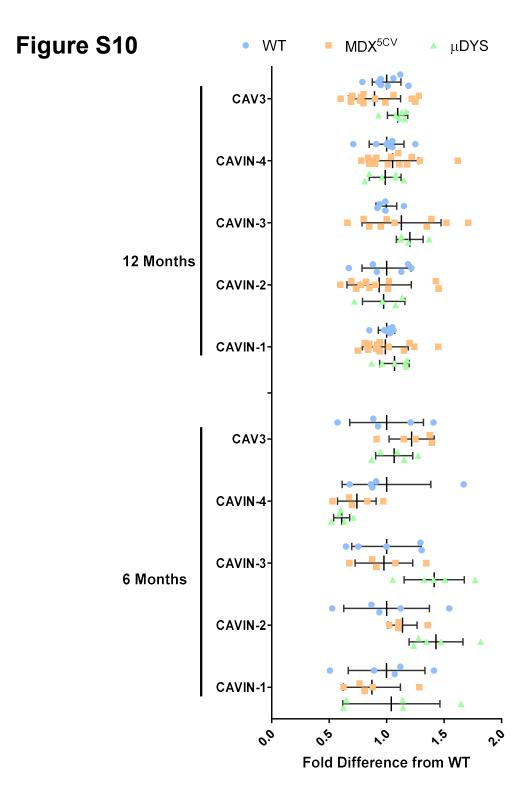


Figure S10: Expression levels of cavins and caveolin 3 in the heart of 6 and 12 months old wild-type,  $mdx^{5cv}$  and  $\mu$ DYS- $mdx^{5cv}$  mice. Protein expression levels were quantified from western blots of total cardiac lysates probed with antibodies to cavins and caveolin-3 (Cav3) Protein levels were normalized to GAPDH probed on the same membrane. Data (mean  $\pm$  standard deviations are expressed as fold differences relative to expression levels in wild-type lysates. No significant differences were found (one-way ANOVA).

Figure S11: Disruption of membrane localization of cavin-4 and cavin-2 in cardiomyocytes from  $mdx^{5cv}$  and  $\mu$ DYS- $mdx^{5cv}$  mice. A. Representative high magnification pictures of tissue sections from of 6 months old wild-type (WT),  $mdx^{5cv}$  and  $\mu$ DYS- $mdx^{5cv}$  ( $\mu$ DYS) mouse hearts double labelled with antibodies to cavin-2 (green) and laminin- $\alpha$ 2 (red). Bright cavin-2 staining is observed in capillaries examples of which are indicated by asterisks. Arrows point to a cardiomyocyte membrane that is not adjacent to a capillary. Scale bar: 10μm. **B.** Representative low magnification micrographs of tissue sections of 6 months old wild-type (WT),  $mdx^{5cv}$  and  $\mu$ DYS- $mdx^{5cv}$  ( $\mu$ DYS) mouse hearts double labelled with antibodies to cavin-4 (green) and dystrophin (red). Scale bar: 50μm.

Figure S11

